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RAW SEQUENCE LISTING

DATE: 04/08/2002

PATENT APPLICATION: US/10/010,050A

TIME: 15:11:42

Input Set : A:\97-38C1Sequence Listing.txt

Output Set: N:\CRF3\04082002\J010050A.raw

P.6

4 <110> APPLICANT: Sheppard, Paul O.
 5 Gilbertson, Debra G.
 7 <120> TITLE OF INVENTION: SECRETED PROTEINS ENCODED BY HUMAN
 8 CHROMOSOME 13
 11 <130> FILE REFERENCE: 97-38C1
 13 <140> CURRENT APPLICATION NUMBER: US 10/010,050A
 C--> 14 <141> CURRENT FILING DATE: 2002-03-26
 16 <150> PRIOR APPLICATION NUMBER: US 09/122,383
 17 <151> PRIOR FILING DATE: 1998-07-24
 19 <150> PRIOR APPLICATION NUMBER: US 60/053,613
 20 <151> PRIOR FILING DATE: 1997-07-24
 22 <160> NUMBER OF SEQ ID NOS: 19
 24 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 26 <210> SEQ ID NO: 1
 27 <211> LENGTH: 1486
 28 <212> TYPE: DNA
 29 <213> ORGANISM: Homo sapien
 31 <220> FEATURE:
 32 <221> NAME/KEY: CDS
 33 <222> LOCATION: (47)...(1084)
 35 <400> SEQUENCE: 1
 36 gaattcggca cgagggcagg aggtagacac ggcacagggc gccgag atg cgg cgg 55
 37 Met Arg Arg
 38 1
 40 ggc gcg ggc gcg gct cgg gga cgc gct tcc tgg tgc tgg gcg ctg gcg 103
 41 Gly Ala Gly Ala Ala Arg Gly Arg Ala Ser Trp Cys Trp Ala Leu Ala
 42 5 10 15
 44 ctg ctt tgg ctc gcg gtg gtt ccg ggc tgg tcc cgg gtc tcg ggc atc 151
 45 Leu Leu Trp Leu Ala Val Val Pro Gly Trp Ser Arg Val Ser Gly Ile
 46 20 25 30 35
 48 ccc tcc cgg cgc cac tgg ccg gtg ccc tac aag cgc ttt gac ttc cgt 199
 49 Pro Ser Arg Arg His Trp Pro Val Pro Tyr Lys Arg Phe Asp Phe Arg
 50 40 45 50
 52 cca aaa cct gat cct tat tgt caa gct aag tat act ttc tgt cca act 247
 53 Pro Lys Pro Asp Pro Tyr Cys Gln Ala Lys Tyr Thr Phe Cys Pro Thr
 54 55 60 65
 56 ggc tca cct atc cca gtt atg gag ggt gat gat gac att gaa gtt ttt 295
 57 Gly Ser Pro Ile Pro Val Met Glu Gly Asp Asp Asp Ile Glu Val Phe
 58 70 75 80
 60 cga tta caa gcc cca gta tgg gaa ttt aaa tat gga gac ctc ctg gga 343
 61 Arg Leu Gln Ala Pro Val Trp Glu Phe Lys Tyr Gly Asp Leu Leu Gly
 62 85 90 95
 64 cac ttg aaa att atg cat gat gcc att gga ttc aga agt aca tta act 391

ENTERED

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65 His Leu Lys Ile Met His Asp Ala Ile Gly Phe Arg Ser Thr Leu Thr
66 100 105 110 115
68 ggc aag aac tac aca atg gaa tgg tat gaa ctt ttc caa ctt ggc aac 439
69 Gly Lys Asn Tyr Thr Met Glu Trp Tyr Glu Leu Phe Gln Leu Gly Asn
70 120 125 130
72 tgt aca ttt ccc cat ctc cga cct gaa atg gat gcc cct ttc tgg tgt 487
73 Cys Thr Phe Pro His Leu Arg Pro Glu Met Asp Ala Pro Phe Trp Cys
74 135 140 145
76 aat caa ggc gct gcc tgc ttt ttt gag gga att gat gat gtt cac tgg 535
77 Asn Gln Gly Ala Ala Cys Phe Phe Glu Gly Ile Asp Asp Val His Trp
78 150 155 160
80 aag gaa aat ggg aca tta gtt caa gta gca act ata tca gga aac atg 583
81 Lys Glu Asn Gly Thr Leu Val Gln Val Ala Thr Ile Ser Gly Asn Met
82 165 170 175
84 ttc aac caa atg gca aag tgg gtg aaa cag gac aat gaa aca gga att 631
85 Phe Asn Gln Met Ala Lys Trp Val Lys Gln Asp Asn Glu Thr Gly Ile
86 180 185 190 195
88 tat tat gag aca tgg aat gta aaa gcc agc cca gaa aag ggg gca gag 679
89 Tyr Tyr Glu Thr Trp Asn Val Lys Ala Ser Pro Glu Lys Gly Ala Glu
90 200 205 210
92 aca tgg ttt gat tcc tac gac tgt tcc aaa ttt gtg tta agg acc ttt 727
93 Thr Trp Phe Asp Ser Tyr Asp Cys Ser Lys Phe Val Leu Arg Thr Phe
94 215 220 225
96 aac aag ttg gct gaa ttt gga gca gag ttc aag aac ata gaa acc aac 775
97 Asn Lys Leu Ala Glu Phe Gly Ala Glu Phe Lys Asn Ile Glu Thr Asn
98 230 235 240
100 tat aca aga ata ttt ctt tac agt gga gaa cct act tat ctg gga aat 823
101 Tyr Thr Arg Ile Phe Leu Tyr Ser Gly Glu Pro Thr Tyr Leu Gly Asn
102 245 250 255
104 gaa aca tct gtt ttt ggg cca aca gga aac aag act ctt ggt tta gcc 871
105 Glu Thr Ser Val Phe Gly Pro Thr Gly Asn Lys Thr Leu Gly Leu Ala
106 260 265 270 275
108 ata aaa aga ttt tat tac ccc ttc aaa cca cat ttg cca act aaa gaa 919
109 Ile Lys Arg Phe Tyr Tyr Pro Phe Lys Pro His Leu Pro Thr Lys Glu
110 280 285 290
112 ttt ctg ttg agt ctc ttg caa att ttt gat gca gtg att gtg cac aaa 967
113 Phe Leu Leu Ser Leu Leu Gln Ile Phe Asp Ala Val Ile Val His Lys
114 295 300 305
116 cag ttc tat ttg ttt tat aat ttt gaa tat tgg ttt tta cct atg aaa 1015
117 Gln Phe Tyr Leu Phe Tyr Asn Phe Glu Tyr Trp Phe Leu Pro Met Lys
118 310 315 320
120 ttc cct ttt att aaa ata aca tat gaa gaa atc cct tta cct atc aga 1063
121 Phe Pro Phe Ile Lys Ile Thr Tyr Glu Glu Ile Pro Leu Pro Ile Arg
122 325 330 335
124 aac aaa aca ctc tct ggt tta taaaacacct taattctact gctctttttt 1114
125 Asn Lys Thr Leu Ser Gly Leu
126 340 345
128 tctccaatca ccagcatctg tttttcaggg ggtgatttta cttttgtgaa ttccttagcc 1174
129 tttcttcctt ggtgcataaa gttaaaatgc acatcagcag aattgctgca tattaacatc 1234

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RAW SEQUENCE LISTING

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Input Set : A:\97-38C1Sequence Listing.txt

Output Set: N:\CRF3\04082002\J010050A.raw

```

130 tcaggactct tctcttgtaa agaagctgaa attcgtacta tattggccaa agtgagcgag 1294
131 ttaggtgata ttggtttcaa tttccgagcc tttgttaata tggagaatta tggttcatat 1354
132 cagttatgta ggacctttgg acccaggggc ctacagatag atatggtgtg cccagatttt 1414
133 aaaaatacct tcaaaaataa aaaatacatt cagtgcacaa aaaaaaaaaa aaaaaatagc 1474
134 ggccgcctcg ag 1486
136 <210> SEQ ID NO: 2
137 <211> LENGTH: 346
138 <212> TYPE: PRT
139 <213> ORGANISM: Homo sapien
141 <400> SEQUENCE: 2
142 Met Arg Arg Gly Ala Gly Ala Ala Arg Gly Arg Ala Ser Trp Cys Trp
143 1 5 10 15
144 Ala Leu Ala Leu Leu Trp Leu Ala Val Val Pro Gly Trp Ser Arg Val
145 20 25 30
146 Ser Gly Ile Pro Ser Arg Arg His Trp Pro Val Pro Tyr Lys Arg Phe
147 35 40 45
148 Asp Phe Arg Pro Lys Pro Asp Pro Tyr Cys Gln Ala Lys Tyr Thr Phe
149 50 55 60
150 Cys Pro Thr Gly Ser Pro Ile Pro Val Met Glu Gly Asp Asp Asp Ile
151 65 70 75 80
152 Glu Val Phe Arg Leu Gln Ala Pro Val Trp Glu Phe Lys Tyr Gly Asp
153 85 90 95
154 Leu Leu Gly His Leu Lys Ile Met His Asp Ala Ile Gly Phe Arg Ser
155 100 105 110
156 Thr Leu Thr Gly Lys Asn Tyr Thr Met Glu Trp Tyr Glu Leu Phe Gln
157 115 120 125
158 Leu Gly Asn Cys Thr Phe Pro His Leu Arg Pro Glu Met Asp Ala Pro
159 130 135 140
160 Phe Trp Cys Asn Gln Gly Ala Ala Cys Phe Phe Glu Gly Ile Asp Asp
161 145 150 155 160
162 Val His Trp Lys Glu Asn Gly Thr Leu Val Gln Val Ala Thr Ile Ser
163 165 170 175
164 Gly Asn Met Phe Asn Gln Met Ala Lys Trp Val Lys Gln Asp Asn Glu
165 180 185 190
166 Thr Gly Ile Tyr Tyr Glu Thr Trp Asn Val Lys Ala Ser Pro Glu Lys
167 195 200 205
168 Gly Ala Glu Thr Trp Phe Asp Ser Tyr Asp Cys Ser Lys Phe Val Leu
169 210 215 220
170 Arg Thr Phe Asn Lys Leu Ala Glu Phe Gly Ala Glu Phe Lys Asn Ile
171 225 230 235 240
172 Glu Thr Asn Tyr Thr Arg Ile Phe Leu Tyr Ser Gly Glu Pro Thr Tyr
173 245 250 255
174 Leu Gly Asn Glu Thr Ser Val Phe Gly Pro Thr Gly Asn Lys Thr Leu
175 260 265 270
176 Gly Leu Ala Ile Lys Arg Phe Tyr Tyr Pro Phe Lys Pro His Leu Pro
177 275 280 285
178 Thr Lys Glu Phe Leu Leu Ser Leu Leu Gln Ile Phe Asp Ala Val Ile
179 290 295 300
180 Val His Lys Lys Gln Phe Tyr Leu Phe Tyr Asn Phe Glu Tyr Trp Phe Leu

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RAW SEQUENCE LISTING

DATE: 04/08/2002

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TIME: 15:11:42

Input Set : A:\97-38C1Sequence Listing.txt

Output Set: N:\CRF3\04082002\J010050A.raw

```

181 305          310          315          320
182 Pro Met Lys Phe Pro Phe Ile Lys Ile Thr Tyr Glu Glu Ile Pro Leu
183          325          330          335
184 Pro Ile Arg Asn Lys Thr Leu Ser Gly Leu
185          340          345
188 <210> SEQ ID NO: 3
189 <211> LENGTH: 18
190 <212> TYPE: DNA
191 <213> ORGANISM: Artificial Sequence
193 <220> FEATURE:
194 <223> OTHER INFORMATION: Oligonucleotide ZC976
196 <400> SEQUENCE: 3
197 cgttgtaaaa cgacggcc
199 <210> SEQ ID NO: 4
200 <211> LENGTH: 17
201 <212> TYPE: DNA
202 <213> ORGANISM: Artificial Sequence
204 <220> FEATURE:
205 <223> OTHER INFORMATION: Oligonucleotide ZC447
207 <400> SEQUENCE: 4
208 taacaatttc acacagg
210 <210> SEQ ID NO: 5
211 <211> LENGTH: 20
212 <212> TYPE: DNA
213 <213> ORGANISM: Artificial Sequence
215 <220> FEATURE:
216 <223> OTHER INFORMATION: Oligonucleotide ZC14487
218 <400> SEQUENCE: 5
219 gacttcgctc caaaacctga
221 <210> SEQ ID NO: 6
222 <211> LENGTH: 20
223 <212> TYPE: DNA
224 <213> ORGANISM: Artificial Sequence
226 <220> FEATURE:
227 <223> OTHER INFORMATION: Oligonucleotide ZC14716
229 <400> SEQUENCE: 6
230 aggggcatcc atttcaggtg
232 <210> SEQ ID NO: 7
233 <211> LENGTH: 20
234 <212> TYPE: DNA
235 <213> ORGANISM: Artificial Sequence
237 <220> FEATURE:
238 <223> OTHER INFORMATION: Oligonucleotide ZC14712
240 <400> SEQUENCE: 7
241 atggctaaac caagagtctt
243 <210> SEQ ID NO: 8
244 <211> LENGTH: 20
245 <212> TYPE: DNA
246 <213> ORGANISM: Artificial Sequence

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RAW SEQUENCE LISTING

DATE: 04/08/2002

PATENT APPLICATION: US/10/010,050A

TIME: 15:11:42

Input Set : A:\97-38C1Sequence Listing.txt

Output Set: N:\CRF3\04082002\J010050A.raw

```

248 <220> FEATURE:
249 <223> OTHER INFORMATION: Oligonucleotide ZC14710
251 <400> SEQUENCE: 8
252 ggggtgaaaca ggacaatgaa      20
254 <210> SEQ ID NO: 9
255 <211> LENGTH: 20
256 <212> TYPE: DNA
257 <213> ORGANISM: Artificial Sequence
259 <220> FEATURE:
260 <223> OTHER INFORMATION: Oligonucleotide ZC14488
262 <400> SEQUENCE: 9
263 ttatgcacca aggaagaaaag      20
265 <210> SEQ ID NO: 10
266 <211> LENGTH: 20
267 <212> TYPE: DNA
268 <213> ORGANISM: Artificial Sequence
270 <220> FEATURE:
271 <223> OTHER INFORMATION: Oligonucleotide ZC14711
273 <400> SEQUENCE: 10
274 ttttctccaa tcaccagcat      20
276 <210> SEQ ID NO: 11
277 <211> LENGTH: 18
278 <212> TYPE: DNA
279 <213> ORGANISM: Artificial Sequence
281 <220> FEATURE:
282 <223> OTHER INFORMATION: Oligonucleotide ZC14430
284 <400> SEQUENCE: 11
285 gtacatttcc ccatctcc      18
287 <210> SEQ ID NO: 12
288 <211> LENGTH: 18
289 <212> TYPE: DNA
290 <213> ORGANISM: Artificial Sequence
292 <220> FEATURE:
293 <223> OTHER INFORMATION: Oligonucleotide ZC14431
295 <400> SEQUENCE: 12
296 ccattttcct tccagtga      18
298 <210> SEQ ID NO: 13
299 <211> LENGTH: 1038
300 <212> TYPE: DNA
301 <213> ORGANISM: Artificial Sequence
303 <220> FEATURE:
304 <223> OTHER INFORMATION: Degenerate nucleotide sequence encoding zsig46
305     polypeptide of SEQ ID NO:2
W--> 307 <221> NAME/KEY: misc_feature
308 <222> LOCATION: (1)...(1038)
309 <223> OTHER INFORMATION: n = A,T,C or G
W--> 311 <400> 13
W--> 312 atgmgngmgng gngcnggngc ngcnmgnggn mgngcnwsnt ggtgytgggc nytngcnytn      60
W--> 313 ytntggytng cngtngtncc nggntggwsn mgngtnwsng gnathccnws nmgnmgncay      120

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 04/08/2002
PATENT APPLICATION: US/10/010,050A TIME: 15:11:43

Input Set : A:\97-38C1Sequence Listing.txt
Output Set: N:\CRF3\04082002\J010050A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:13; N Pos. 6,9,12,15,18,21,24,27,30,33,36,39,51,54,57,60,63,69,72,75
Seq#:13; N Pos. 78,81,84,90,93,96,99,102,108,111,114,117,126,129,132,141
Seq#:13; N Pos. 153,156,162,168,180,189,198,201,204,207,210,216,219,228,246
Seq#:13; N Pos. 252,255,261,264,267,285,291,294,297,303,321,327,333,336,339
Seq#:13; N Pos. 342,345,348,360,378,387,390,399,405,411,414,417,429,432,450
Seq#:13; N Pos. 453,456,471,483,501,504,507,510,516,519,522,528,531,552,561
Seq#:13; N Pos. 579,582,597,606,612,615,618,627,630,636,648,660,669,672,675
Seq#:13; N Pos. 678,690,693,702,705,726,735,738,747,753,756,762,765,771,774
Seq#:13; N Pos. 783,786,789,795,798,801,804,813,816,819,822,825,834,846,855
Seq#:13; N Pos. 861,864,867,879,882,885,888,891,906,909,915,933,960,963,975
Seq#:13; N Pos. 990,1005,1008,1011,1017,1026,1029,1032,1035,1038